## STIC-Biotech/ChemLib

From:

Chernyshev, Olga

Sent:

Monday, March 08, 2004 1:16 PM

To: Subject: STIC-Biotech/ChemLib sequence search request

US case 09/905,075. Interference search for SEQ ID NO: 2. Please print out a summary table of direct 100% hits. Thank you very much!

Olga N. Chernyshev AU1646 REM 4E84 20870 mail box 4D70

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Searcher: FORT
Phone:
Location:
Date Picked Up: 3///2
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Date Completed: / 3//0
Searcher Prep/Review:///
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Online time:

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other

/ENDOR/COST (w	here applic.)
STN:	
DIALOG:	
Questel/Orbit:	
DRLink:	
Lexis/Nexis:	
Sequence Sys.:/	
WWW/Internet:_	<u> </u>
Other (specify):	

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Searcher:
Phone:
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Date Picked Up:
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TYPE OF SEARCH:	
NA Sequences:	
AA Sequences:	
Structures:	
Bibliographic:	_
Litigation:	
Full text:	
Patent Family:	_
Other:	

/ENDOR/COST (where applic.) STN:
DIALOG:
Questel/Orbit:
DRLink:
Lexis/Nexis:
Sequence Sys.:
WWW/Internet:
Other (checify):

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                        OM protein - protein search, using sw model
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US-09-905-075-2 2005 1 WRLPRRAALGLLPLLLLLPP......AEAEATEGESPTQLPSREDL 353 389414 seqs, 51625971 residues BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table: Perfect score: Sequence: Title:

389414 Total number of hits satisfying chosen parameters:

Searched:

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

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/cgn2\_6/ptodata/2/iaa/PCTUS\_COMB.pep:\*
/cgn2\_6/ptodata/2/iaa/PCTUS\_COMB.pep:\* Issued Patents AA:\* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

		ð			SOUTHERE	
Result No.	Score	Query Match	Length	DB	αı	Description
	2005	100.0	353	4	US-09-907-794A-2	Sequence 2, Appli
7	2005	100.0	353	4	US-09-905-125A-2	7
m	2005	100.0	353	4	US-09-902-775A-2	Sequence 2, Appli
4	971.5	48.5	420	4	US-09-907-794A-109	100
Ŋ	971.5	٠,	420	4	-60	109,
9	971.5		420	4	US-09-902-775A-109	10
7	463.5	23.1	242	4	39-312	393,
80	256	12.8	77	4	US-09-621-976-4010	4010,
6	RJ.	12.7	1253	٣	US-08-479-722B-4	Sequence 4, Appli
10	254.5	12.7	652	7	US-08-751-305-2	7
11	250	12.5	1833	m	US-08-479-722B-2	2,7
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13	246.5	12.3	1251	Ŋ	PCT-US95-02251-3	3, 1
14	246.5	12.3	1252	Н	- 1	'n
15	246.5	12.3	1252	~	US-08-316-650-3	'n
16	240.5	12.0		9	5177197-30	ίς
17	239	11.9	919	Н	US-08-282-141-4	Sequence 4, Appli
18	239	11.9	979	Н	US-08-435-434-3	e,
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22	239	11.9		3	$\tilde{\sim}$	e,
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26	238	11.9	509	4	US-09-907-794A-315	Sequence 315, App
27	238	11.9		4	US-09-905-125A-315	31

315 233, 223, 221, 22, 7, 12, 15,	Sequence 2, Appli Sequence 2, Appli Patent No. 5258288 Sequence 5, Appli
US-09-902-775A-315 US-08-525-940-23 US-08-976-838-23 US-08-976-838-21 US-08-976-838-21 US-08-976-838-18 US-08-976-838-18 US-09-214-555B-7 US-09-341-461-2 US-09-341-461-2 US-08-525-940-15 US-08-938-15 US-08-938-15 US-08-96-96-15	US-08-436-804-2 US-08-267-387-2 5258288-4 US-08-882-046-5
4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	3611
509 799 881 881 915 915 915 3623 288 288 635 635	635 635 652 1219
	11.3 11.2 11.2
238 236 236 236 236 236 236 230 230 227.5	227.5 227.5 224 224
90010000000000000000000000000000000000	4 4 4 4 2 & 4 2

## ALIGNMENTS

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APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
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CURRENT APPLICATION NUMBER: US/09/907,794A
CURRENT FILING DATE: 2001-07-17
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR PRILING DATE: 1999-07-07
PRIOR PELING DATE: 1999-07-07
PRIOR PELING DATE: 1999-07-07
PRIOR PELING DATE: 1999-07-07
PRIOR PLING DATE: 1999-07-26
PRIOR PLING DATE: 1999-07-26
PRIOR PLING DATE: 1999-07-28
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PRIOR PLING DATE: 1999-07-28
PRIOR PLING DATE: 1999-09-08
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PRIOR PRING DATE: 1999-09-08
PRIOR PRING DATE: 1999-09-13
PRIOR PRILING DATE: 1999-09-13
PRIOR PRILING DATE: 1999-09-15
PRIOR PRILING DATE: 1999-09-15
PRIOR PRILING DATE: 1999-09-15
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Grimaldi, Christopher J.
Grimaldi, Christopher J.
Hillan, Kenneth, J.
Kljavin, Ivar J.
Mather, Jennie P.
Pan, James
Paoni, Nicholas F.
Roy, Margaret Ann
Stewart, Timothy A.
Tumas, Daniel
                                      Sequence 2, Application US/09907794A Patent No. 6635468 GENERAL INFORMATION:
                                                                                                                                                                                                                             Ferrara, Napoleone
Filvaroff, Ellen
                                                                                                                                                                                                                                                                                                                         Gerber, Hanspeter
Gerritsen, Mary E.
                                                                                                            APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
                                                                                                                                                                                    Desnoyers, Luc
Eaton, Dan L.
                                                                                                                                                                                                                                                                              Fong, Sherman
Gao, Wei-Qiang
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                     US-09-907-794A-2
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RESULT 1
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us-09-905-075-2.rai

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VKTLKVCCSPGTYGPDCLACQGGSQRPCSGNGHCSGDGSRQGDGSCRCHMGYQGPLCTDC 180
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Pred. No. 2.6e-150;
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TITLE OF INVENTION: Acids Encoding the Same
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/145,048
PRIOR FILING DATE: 1999-07-26
PRIOR FILING DATE: 1999-07-26
PRIOR FILING DATE: 1999-07-26
PRIOR FILING DATE: 1999-07-26
PRIOR PELICATION NUMBER: US 60/146,222
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR PELING DATE: 1999-09-08
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PRIOR PELING DATE: 1999-09-13
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PRIOR PELING DATE: 1999-12-07
PRIOR PELING DATE: 1999-12-06
PRIOR PELING DATE: 1999-12-06
PRIOR PELING DATE: 1999-12-06
PRIOR PELING DATE: 1999-12-07
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CURRENT APPLICATION NUMBER: US/09/905,125A
CURRENT FILING DATE: 2001-07-12
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PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2007-07-12
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           Grimaldi, Christopher
                                 Gurney, Austin L.
Hillan, Kenneth, J.
Kljavin, Ivar J.
Mather, Jennie P.
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Paoni, Nicholas F.
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Best Local Similarity 100.0
Matches 353; Conservative
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TYPE: PRT
CRGANISM: Homo sapiens
US-09-905-125A-2
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APPLICANT:
APPLICANT:
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100.0%; Pred. No. 2.6e-150;
iive 0; Mismatches 0;
                                             PRIOR PILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR PLING DATE: 1999-11-29
PRIOR PLING DATE: 1999-11-20
PRIOR PILING DATE: 1999-12-02
PRIOR PLING DATE: 1999-12-04
PRIOR PLING DATE: 1999-12-06
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PRIOR PLING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR PLING DATE: 2000-01-05
PRIOR PLING DATE: 2000-01-05
FILING DATE: 1999-09-15
APPLICATION NUMBER: PCT/US99/23089
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Patent No. 6664376
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Gao, Wei-Qiang
Gerber, Hanspeter
Gerritsen, Mary E.
Goddard, A.
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Filvaroff, Ellen
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Ashkenazi, Avi
Botstein, David
Desnoyers, Luc
Eaton, Dan L.
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Best Local Simi:
Matches 353; (
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; ORGANISM: Hon
US-09-907-794A-2
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Paoni, Nicholas F.
Roy, Margaret Ann
Stewart, Timothy A
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US-09-907-794A-109
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US-09-902-775A-2
                                                                                                                                                                                                                              LENGTH: 353
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APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT PILICATION NUMBER: US/09/902,775A
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR FILING DATE: 1999-07-28
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MDGYFSSLRNETHSICTACDESCKTCSGLTNRDCGECEVGWVLDEGACVDVDECAAEPPP 240
                            MDGYFSSLRNETHSICTACDESCKTCSGLTNRDCGECEVGWVLDEGACVDVDECAAEPPP
                                                                                CSAAQFCKNANGSYTCEECDSSCVGCTGEGPGNCKECISGYAREHGQCADVDECSLAEKT
                                                                                                                     241 CSAAQFCKNANGSYTCEECDSSCVGCTGEGPGNCKECISGYAREHGQCADVDECSLAEKT
                                                                                                                                                                 CVRKNENCYNTPGSYVCVCPDGFEETEDACVPPARARATEGESPTQLPSREDL 353
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PRIOR FILING DATE: 1999-09-13
PRIOR PELLING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR APPLICATION NUMBER: PCT/US99/21647
PRIOR FILING DATE: 1999-09-15
PRIOR FILING DATE: 1999-09-15
PRIOR PLING DATE: 1999-10-05
PRIOR PLING DATE: 1999-11-09
PRIOR PLING DATE: 1999-11-29
PRIOR PLING DATE: 1999-11-29
PRIOR PLING DATE: 1999-11-29
PRIOR PLING DATE: 1999-11-30
PRIOR PLING DATE: 1999-11-29
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PRIOR PLING DATE: 1999-11-29
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Grimaldi, Christopher J.
Gurney, Austin L.
Hillan, Kenneth, J.
                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/09902775A Patent No. 6686451 GENERAL INFORMATION:
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Paoni, Nicholas F.
Roy, Margaret Ann
Stewart, Timothy A.
Tumas, Daniel
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Filvaroff, Ellen
Fong, Sherman
Gao, Wei-Qiang
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Goddard, A.
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Mather, Jennie P.
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Ashkenazi, Avi
Botstein, David
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Eaton, Dan L.
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61 EEKTLSKYESSEIRLLEILEGLCESSDFECNQMLEAQEEHLEAWWLQLKSEYPDLFEWFC 120
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100.0%; Score 2005; DB 4;
Best Local Similarity 100.0%; Pred. No. 2.6e-150;
Matches 353; Conservative 0; Mismatches 0;
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FLILING DATE: 1999-12-02
PRIOR PLILING DATE: 1999-12-02
PRIOR PLILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR PLILING DATE: 1999-12-20
PRIOR PLILING DATE: 1999-12-20
PRIOR PLILING DATE: 1999-12-20
PRIOR FLILING DATE: 2000-01-05
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Grimaldi, Christopher J.
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Filvaroff, Ellen
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Gao, Wei-Qiang
Gerber, Hanspeter
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Mather, Jennie P.
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APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 423
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Homo sapiens
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GENERAL INFORMATION:
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APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Scretced and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
                                 311 -TEVCPGENKQCENTEGGYRCICAEGYKQMEGICVKEQIPESAGFFSEMTEDE 362
297 AEKTCVRKNENCYNTPGSYVCVCPDGFEETEDACVP---PAEA
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PRIOR PLING DATE: 1999-07-07
PRIOR PLING DATE: 1999-07-07
PRIOR PLING DATE: 1999-07-07
PRIOR PLING DATE: 1999-07-06
PRIOR PLING DATE: 1999-07-28
PRIOR PLING DATE: 1999-07-28
PRIOR PELING DATE: 1999-09-09
PRIOR PLING DATE: 1999-09-09
PRIOR PLING DATE: 1999-09-13
PRIOR PLING DATE: 1999-09-13
PRIOR PLING DATE: 1999-09-15
PRIOR PLING DATE: 1999-10-01
PRIOR APPLICATION NUMBER: PCT/US99/28314
PRIOR PLING DATE: 1999-10-16
PRIOR APPLICATION NUMBER: PCT/US99/28314
PRIOR PLING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR PLING DATE: 1999-11-30
PRIOR PLING DATE: 1999-11-20
PRIOR PLING DATE: 1999-11-20
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PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
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PRIOR APPLICATION NUMBER: US 60/143,048
                                                                                                                                                                                       Sequence 109, Application US/09905125A Patent No. 6664376
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Grimaldi, Christopher J.
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Eaton, Dan L.
Ferrara, Napoleone
Filvaroff, Ellen
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Hillan, Kenneth, J.
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Mather, Jennie P.
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Ashkenazi, Avi
Botstein, David
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Gao, Wei-Qiang
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                                                                                                                                                                  JS-09-905-125A-109
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         APPLICANT: Tunas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, Williams, P. Mickey
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acide Encoding the Same
FILE REFERENCE: 10466-14
CURRENT FILING DATE: 2001-07-17
PRIOR PLILING DATE: 2000-02-22
PRIOR PLILING DATE: 2000-02-22
PRIOR PLILING DATE: 1999-07-07
PRIOR PLILING DATE: 1999-07-07
PRIOR PLILING DATE: 1999-07-26
PRIOR PLILING DATE: 1999-09-08
PRIOR PLILING DATE: 1999-09-08
PRIOR PLILING DATE: 1999-09-15
PRIOR PLILING DATE: 1999-10-57
PRIOR PRIOR PLILING DATE: 1999-10-57
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PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 109
LENGTH: 420
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Best Local Similarity 48.2
Matches 170; Conservative
   Tumas, Daniel
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ORGANISM: Homo sapiens
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APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
                                                                                                                                                                               7 AALGLIPILILILP-----PAPEAA--KKPTPCHRCRGLVDKFNQGMVDTAKKNFGGG
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                                                                                                                     Length 420;
                                                                                                                                                   Indels
                                                                                                                        DB 4;
                                                                                                                    48.5%; Score 971.5; DB 4;
48.2%; Pred. No. 7e-69;
ative 48; Mismatches 116;
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CURRENT APPLICATION NUMBER: US/09/902,775A
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: PCT/US00/04414
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Sequence 109, Application US/09902775A
Patent No. 6686451
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Grimaldi, Christopher J.
Gurney, Austin L.
Hillan, Kenneth, J.
Kljavin, Ivar J.
Mather, Jennie P.
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Stewart, Timothy A.
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Gerber, Hanspeter
Gerritsen, Mary E.
Goddard, A.
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Ferrara, Napoleone
Filvaroff, Ellen
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   2000-01-05
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Abhkenazi, Avi
Botstein, David
Desnoyers, Luc
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Stewart, ....
Stewart, ....
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PRIOR FILING DATE: 2000-0:
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 109
LENGTH: 420
                                                                         , ORGANISM: Homo sapiens
US-09-905-125A-109
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                                                                                                                  Query Match
Best Local Similarity
Matches 170; Conserv
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APPLICANT:
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57 NTAWEEKTLSKYESSEIRLLEILEGLCESSDFECNOMLEAQEEHLEAWWLQLKSEYPDLF 116
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PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-07
PRIOR FILING DATE: 1999-07-26
PRIOR PELING DATE: 1999-07-26
PRIOR PELING DATE: 1999-07-28
PRIOR PELING DATE: 1999-07-28
PRIOR PELING DATE: 1999-09-08
PRIOR PELING DATE: 1999-09-08
PRIOR PELING DATE: 1999-09-13
PRIOR PELING DATE: 1999-09-13
PRIOR PELING DATE: 1999-09-13
PRIOR PELING DATE: 1999-09-15
PRIOR PELING DATE: 1999-10-05
PRIOR PELING DATE: 1999-11-20
PRIOR PELING DATE: 1999-11-20
PRIOR PELING DATE: 1999-11-30
PRIOR PELING DATE: 1999-11-30
PRIOR PELING DATE: 1999-11-30
PRIOR PELING DATE: 1999-11-20
PRIOR PELING DATE: 1999-12-02
PRIOR PELING DATE: 1999-12-02
PRIOR PELING DATE: 1999-12-16
PRIOR PELING DATE: 1999-12-20
PRIOR PELING DATE: 1999-12-30
PRIOR PELING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 109
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; Patent No. 6573095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  al Similarity 48.2
170; Conservative
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US-09-902-775A-109
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Best Local &
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COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                             Sequence 4, Application US/08479722B Patent, No. 6074840 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     : 1253 amino acids
amino acid
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
                                                                                                                                                                                                                       STREET: 7676 1
CITY: Houston
                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                           STATE: Texas
COUNTRY: USA
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             US-08-479-722B-4
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                                                                                                                                                                                                                                                                                                                                                                      DB 4; Length 242;
APPLICANT: Watson, James D.
APPLICANT: Watson, James D.
APPLICANT: Steman, Matthew
APPLICANT: Sleeman, Matthew
APPLICANT: Chrust, Rene
APPLICANT: Murison, James G.
APPLICANT: Munble, Krishanand D.
TITLE OF INVENTION: Compositions Isolated from Skin Cells
TITLE OF INVENTION: Compositions Isolated from Skin Cells
TITLE OF INVENTION: Campositions (or Their Use)
FILE REPRESENCE: 11000.1011c2
CURRENT FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 425
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 393
LENGTH: 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 77;
                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4010, Application US/09621976

Patent No. 6639063

GENERAL INFORMATION:
APPLICANT: Unuas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Glocatdano, J.Y.
ITILE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976

CURRENT FILING DATE: 2000-07-21

NUMBER OF SEQ ID NOS: 19335
                                                                                                                                                                                                                                                                                                                                                                  23.1%; Score 463.5; DB 4;
44.6%; Pred. No. 3.7e-29;
tive 18; Mismatches 76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12.8%; Score 256; DB 4;
94.1%; Pred. No. 2.3e-13;
iive 2; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                        83; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 94.1
Matches 48; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EATEGE 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            179 EMTEDE 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; LOCATION: -24..-1
US-09-621-976-4010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: SIGNAL
                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Mouse
                                                                                                                                                                                                                                                                                                                               US-09-312-283C-393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-621-976-4010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 4010
LENGTH: 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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606 GPGKGICMNTGGSY-----NCHCNRGYRLHVGAGGRSCVDLNECAKPHLCGDGGFCINFP 660
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 PAPEAAKKPTPCHRCRGLVDKFNQGMVDTAKKNFGGGNTAWEEKTLSKYESSEIRLLEIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                518 PYPELISRPSP-----PTFHRFLPDLPP----SRSAVEIAPTQVTETDECRLNONI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   165 ---SCRCHMGY----QGPLCTD-----CMDGYFSSLRNETHSI-CTACDESCKTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ----GACRDVNEC-SEGIPCSPG-WCENLPGSYRCTCAQGIRTRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   80 --EGLC--ESSDF--ECNOMLEAQEEHLEAWWLQLKSEYPDLFEWFCVKTLKVCCSPGTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -----GDG----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         661 GHYKCNCYPGYRLKASRPPICEDIDECRDPSTCPDG---KCENKPGSFKCIACQPGYRSQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1253;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      207 SGLTNRDCGECEVGWVLDEGACVDVDECAAEPPFCSAAQFCKNANGSYTC-
APPLICANT: Bonadio, Jeffrey
APPLICANT: Yin, Wushan
TITLE OF INVENTION: LATENT TGF( BINDING PROTEIN (LTBP)
TITLE OF INVENTION: LATENT TGF( BINDING PROTEIN (LTBP)
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            134 GPDCLACQ--GGSQRPCSGNGHCS-----GDGSRQ-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/479,722B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
12.7%; Score 255.5; DB 3;
Best Local Similarity 24.0%; Pred. No. 5.3e-12;
Matches 102; Conservative 40; Mismatches 109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 08/199,780
FILING DATE: 18-FEB-1994
ATTOKING DATE: 18-FEB-1994
ATTOKING DATE: 18-FEB-1994
NAME: FUSSEY, Shelley P.M.
REGISTRATION NUMBER: 39,458
REFERENCE/DOCKET NUMBER: 4100.000500/FUS
TELEPHONE: (713) 934-7001
TELEPHONE: (713) 934-7011
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             565 CGHĠQĊVPGPŚDYSCHĊNAGYRSHPQHR-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFCATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US PCT/US95/02251
FILING DATE: 21-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/316,650
FILING DATE: 30-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/199,780
                                                                                                                                                                                            ADDRESSEE: Williams, Morgan & Amerson
STREET: 7676 Hillmont, Suite 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UMBER: US/08/479,722B
07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         718 GG----
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997 VNSPG--SYTCLACEEGYVGQSGSCVDVNECLTPGICTHGRCINMEGSFRCSCEPGYEVT 1054
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       128 CSPG-----TYGPDCLACQGGSQRPCSGNGHCSGDGSRQGDGSCRCHMGY------QG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      937 QECODIDECEQPGVCSGGRCSNTEGSYHCECDRGYIMVRKGHCQDINECRHPGTCPDGRC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           --SCVG----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ----CAAEP----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12.5%; Score 250; DB 3; Length 1833; 24.8%; Pred. No. 2.2e-11; ative 22; Mismatches 94; Indels 13
                                                                                                                                          Sequence 2, Application US/08479722B
Patent No. 6074840
GENERAL INFORMATION:
APPLICANT: Bonadio, Jeffrey
APPLICANT: Yin, Wushan
TITLE OF INVENTION: LATENT TGF( BINDING PROTEIN (LTBP)
TITLE OF INVENTION: LATENT TGF( BINDING PROTEIN (LTBP)
TITLE OF INVENTION: GENES, COMPOSITIONS AND METHODS:
CORRESPONDENCE ADDRESS:
ADDRESSER: Williams, Morgan & Amerson
STREET: 7676 Hillmont, Suite 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/479.722R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -----PPCSA----AQFCKNANGSYTCEECDS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4100.000500/FUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US PCT/US95/02251
PILING DATE: 21-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/316,650
PILING DATE: 30-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/199,780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>..</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   204 KTCSGLTNRDCGECEVGWVLDEGACVDVDE-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UMBER: US/08/479,722B
07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: FUSSEY, Shelley P.M. REGISTRATION NUMBER: 39,458 REFERENCE/DOCKET NUMBER: 41 TELECOMMUNICATION INFORMATION: (713) 934-7000
325 ----ETEDACVPP 333
                                        421 DGTQCQDVDECVGP 434
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1833 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 18-FEB-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (713) 934-7011
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query,Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: SOFTWARE: Patent:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 0'CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                    STREET: 7676 ]
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                               Texas
: USA
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                                                                                                                                                                                                                                                                                                                                                                                                             STATE: Texal
COUNTRY: USI
ZIP: 77040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                        RESULT 11
US-08-479-722B-2
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             ------EECDSSCV---GCTGEGPGNCK-ECISGY--AREHGQCADVDECSLAEK--- 299
                                                                                                                            |: | : | HINDESYRCLCPLGHRLVGGRKCKKDIDECSQDPGLCLPHACENLQGSYVCVCDEGF 878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             212 RDCGECEVGWVLDEGA--CVDVDECAAEPPCSAAQFCKNANGSYTCEECDSSC-VGCTG 268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     326 YTC-RCPQGYQLDSSQLDCVDVDBC--QDSPC--AQECVNTPGGFRCE-----CWYGYEP 375
                                      759 GRLSCIDVDECEAGKVCQDGICTNTPGSFQCQCLSGYHLSRDRSRCEDIDECDFPAACIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                154 CSGDGSRQGDGS--CRCHMGYQGPLCTDCMDGYFSSLRNETHSICTACDESCKTCSGLTN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    269 EGPGNCKECISGYAREHGQCADVDECSLAEKTCVRKNENCYNTPGSYVCVCPDGFE----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95 EAQEEHLEAWWLQLKSEYPDLFEWFCVKTLKVCCSPGTYGPDCLACQGGSQRPCS-GNGH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12.7%; Score 254.5; DB 2; Length 652; larity 30.7%; Pred. No. 3.1e-12; Conservative 27; Mismatches 88; Indels 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COPETWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/751,305
FILING DATE: 18-NOV-1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell, Jr., John R.
REGISTRATION NUMBER: 31,678
REFERENCE/DOCKET NUMBER: 31,678
REFERENCE/DOCKET NUMBER: 07306/012001
TELECOMMULCATION INFORMATION:
TELECOMMULCATION INFORMATION:
TELECOMMULCATION INFORMATION:
TELECOMMULCATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/08751305
Patent No. 5965439
GENERAL TON:
TYPE TENNER E al., Andrea J.
TITLE OF INVENTION: HOST DEFENSE ENHANCEMENT NUMBER OF SEQUENCES: 33
CORRESPONDENCES:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : 652 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 78; Conserv
                                                                                                 300 -TCVRKNEN-
                                                                                                                                                                                     324 EETED 328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 4220
                                                                                                                                                                                                                                TLTOD 883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              376 GGPG--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: La Jo
STATE: CA
COUNTRY: US
ZIP: 92037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                             RESULT 10
US-08-751-305-2
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997 VNSPG--SYTCLACEEGYVGQSGSCVDVNECLIPGICTHGRCINMEGSFRCSCEPGYEVT 1054
                                                                                                                                                                                                                                                            266 ----CTGE-GPGNCKECISGYARE--HGQCADVDECSLAEKTCVRKNENCYNTPGSYVCV 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 80 --EGLC--ESSDF--ECNOMLEAQEEHLEAWWLQLKSEYPDLFEWFCVKTLKVCCSPGTY 133
                                                                                                                                                                   1055 PDKKGCRDVDECASRASCPIGLCLNTEGSFTCSACQSGYWVNEDGTACEDLDECAFPGVC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        METHODS AND COMPOSITIONS FOR STIMULATING BONE CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
12.3%; Score 246.5; DB 5; Length 1251;
Best Local Similarity 23.5%; Pred. No. 2.7e-11;
Matches 100; Conservative 41; Mismatches 109; Indels 175;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: United States of America
ZIP: 77210
                                                                                        239 -----PPCSA----AQFCKNANGSYTCEECDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          319 CPDGFEET------EDACVPPAE 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
TITLE OF INVENTION: METHODS AND COMPOSITIONE OF INVENTION: CELLS
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSER: Arnold, White & Durkee
STREET: P.O. Box 4433
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3, Application PC/TUS9502251
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LASTICS:
LASTICS:
TYPE: amino acids
TOPOLOGY: ling
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(713) 789-2679
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TELEEX: 79-0924
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                    1055 PDKKGCRDVDECASRASCPTGLCLNTEGSFTCSACQSGYWVNEDGTACEDLDECAFPGVC 1114
                                                                               ----CTGE-GPGNCKECISGYARE--HGQCADVDECSLAEKTCVRKNENCYNTPGSYVCV 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   883 CSPGYQLHPSQDYCTDDNECM---RNPCEGRGRCV---NSVGSYSCLCYPGYTLVTLGDT 936
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                                                                                                                                                                                                                                                        -----EDACVPPAE 335
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PCT-US95-02251-18
SCHOOLE 18, Application PC/TUS9502251
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: CELLS
ITITE OF SEQUENCES: 18
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee STREET: P.O. Box 4433
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REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UMIC009P---
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: CONCURRENTLY HEREWITH CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/316,650
FILING DATE: 30-SEP-1994
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 08/199,780
FILING DATE: 18-FEB-1994
CLASSIFICATION:
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INFORMATION FOR SEQ ID NO: 18:
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TELEFAX: (713) 789-2679
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Best Local Similarity 24.8<sup>†</sup>
Matches 82, Conservative
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CITY: Houston
STATE: Texas
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718 GG
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US-08-316-650-3
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--YCVDVNE--CEAEPC 604
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                                                                                                                                              660 GHYKCNCYPGYRLKASRPPICEDIDECRDPSTCPDG---KCENKPGSFKCIACQPGYRSQ
                                                                                                                                                                                                                                                           257 -----EECDSSCV---GCTGEGPGNCK-ECISGY--AREHGQCADVDECSLAEK---
                                                                                                                                                                                                                                                                              818 GDCINTNGSYRCLCPLGHRLVGGRKCKKDIDECSQDPGLCLPHACENLQGSYVCVCDEGF
                                                                                                         ---SCRCHMGY-----QGPLCTD------CMDGYFSSLRNETHSI-CTACDESCKTC
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                                                                                                                                                                                   207 SGLTNRDCGECEVGWVLDEGACVDVDECAAEPPFCSAAQFCKNANGSYTC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3, Application US/08199780

Patent No. 5763416

GENERAL INFORMATION:
APPLICANT: Bonadio, Jeffrey
APPLICANT: Goldstein, Steven A.
TITLE OF INVENTION: Gene Transfer Into Bone Cells
TITLE OF INVENTION: And Tissues
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: AFROIGH, White & Durkee
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FAPLICATION NUMBER: US/08/199,780
FILING DATE: 18-FEB-1994
CLASSIFICATION: 514
564 CGHGQCVPGPSDYSCHCNAGYRSHPQHR--
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                                    134 GPDCLACQ -- GGSQRPCSGNGHCS--
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UM
TELECOMMUNICATION INFORMATION:
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INFORMATION FOR SEQ ID NO: 3:
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ATTORNEY/AGENT INFORMATION:
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CITY: Houston
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Matches 100; Conserv
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165 ---SCRCHMGY-----QGPLCTD-----CMDGYFSSLRNETHSI-CTACDESCKTC 206
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80 --EGLC--ESSDF--ECNOMLEAQEEHLEAWWLQLKSEYPDLFEWFCVKTLKVCCSPGTY 133
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                                                                                                                                                                                                                                         606 GPGKGICMNTGGSY-----NCHCNRGYRLHVGAGGRSCVDLNECAKPHLCGDGGFCINFP
                                                                                                                                                                                                                                                                                                                                                                                               207 SGLTNRDCGECEVGWVLDEGACVDVDECAAEPPPCSAAQFCKNANGSYTC
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/316,650
FILING DATE: 30-SEP-1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/199,780
FILING DATE: 30-SEP-1994
ATTORNEY/AGENT INFORMATION:
NAME: PARKET, DAVIG L.
NAME: PARKET, DAVIG L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UMIC:008
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lin, Wushan
TITLE OF INVENTION: METHODS AND COMPOSITIONS
TITLE OF INVENTION: POR STIMULATING BONE CELLS
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                          134 GPDCLACQ--GGSQRPCSGNGHCS-----GDGSRQ--
                                                                                                                                      565 СGHGQCVPGPSDYSCHCNAGYRSHPQHR-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3, Application US/08316650
Patent No. 5942496
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roessler, Blake J.
Goldstein, Steven A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Bonadio, Jeffrey
APPLICANT: Roessler, Blake
APPLICANT: Goldstein, Steve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (512) 418-3000
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MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                 565 CGHGQCVPGPSDYSCHCNAGYRSHPQHR-------YCVDVNE--CEAEPC 605
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                                                                                                                                                                                                                                                                                                                                                                        606 GPGKGICMNTGGSY-----NCHCNRGYRLHVGAGGRSCVDLNECAKPHLCGDGGFCINFP 660
                                                                                                                                                                                                                                                                                                                                                                                                             165 ---SCRCHMGY-----QGPLCTD-----CMDGYFSSLRNETHSI-CTACDESCKTC 206
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                                                                                                                                                                                    20 PAPEAAKKPTPCHRCRGLVDKFNQGMVDTAKKNFGGGNTAWEEKTLSKYESSEIRLLEIL 79
                                                                                                                Query Match 12.3%; Score 246.5; DB 2; Length 1252; Best Local Similarity 23.5%; Pred. No. 2.7e-11; Matches 100; Conservative 41; Mismatches 109; Indels 175; Gaps
LENGTH: 1252 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       300 -TCVRKNEN-
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US-08-316-650-3
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Search completed: March 9, 2004, 16:00:05 Job time: 25 secs

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